



SEQUENCE LISTING

Iedhar, Shoukat
Hannigan, Greg
Yee, Arthur

120 INTEGRIN-LINKED KINASE AND ITS USES

130 FINE001C1P4

140 09/323,548

141 2001-03-08

150 09/320,435

151 1999-03-03

150 09/335,706

151 1998-03-05

150 08/955,841

151 1997-10-21

150 08/752,845

151 1996-11-19

150 03/009,074

151 1995-12-21

160 33

170 FastSEQ for Windows Version 4.0

210 1

211 1759

212 DNA

213 Homo sapiens

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acg gag aac gac ctg aac cag ggg gac gat cat ggc ttc tcc ccc ttg	270
cac tgg gcc tgc cga gag ggc cgc tct gct gtg gtt gag atg ttg atc	318
atg cgg ggg gca cgg atc aat gta atg aac cgt ggg gat gac acc ccc	366
ctg cat ctg gca gcc agt cat gga cac cgt gat att gta cag aag cta	414
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ccc ctg cag tat gcc tgt ttt tgg ggc caa gat caa gtg gca gag gac	510
ctg gtg gca aat ggg gcc ctt gtc agc atc tgt aac aag tat gga gag	558
atg cct gtg gac aaa gcc aag gca ccc ctg aga gag ctt ctg cga gag	606
ggg gca gag aac atg ggc cag aat ctg aac cgt att cca tac aag gac	654
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aac aaa cac tct ggc att gac ttc aaa cag ctt aac ttc ctg acg aag	750
ctc tac gag aat cac tct gga gag cta tgg aag ggc cgc tgg cag gcc	798
aat cac att gtc gtg aag gtg ctg aag gtt cga gac tgg agt aca agg	846
aag agc agg gac ttc aat gaa gag tgt ccc cgg ctg agg att ttc cag	894
cat cca aat gtg ctg cca gtg cta ggt gcc tgc cag tct cca cct gct	942

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MAY 14 2003

TECH CENTER 1600/2900

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 aat gta cta cat gaa ggc acc aat ttc gtc gtg gac cag agc cag gct 1038
 gag aag ttt gct ttg gac atg gca agg ggc atg gcc ttc cta cac aca 1086
 cta gag ccc ctc atc cca cga cat gca ctc aat agc cgt agt gta atg 1134
 att gat gag gac atg act gcc cga att agc atg gct gat gtc aag ttc 1182
 ttt ttc caa tgt cct ggt cgc atg tat gca cct gcc tgg gta gcc ccc 1230
 gaa gct ctg cag aag aag cct gaa gac aca aac aga cgc tca gca gac 1278
 atg tgn agt ttt gca gtg ctt ctg tgg gaa ctg gtg aca cgg gag gta 1326
 ccc ttt gct gac ctc tcc aat atg gag att gga atg aag gtc gca ttg 1374
 gaa ggt ctt cgg cct acc atc cca cca ggt att tcc cct cat gtg tgt 1422
 aag ctc atg aag atc tgc atg aat gaa gac cct gca aag cga ccc aaa 1470
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 taggactgga aggtccttgc ctgaactcca gaggtgtcgg gacatggttg ggggaatgca 1572
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 gcttgcacac ttgcacatg gtgtctccca acatgggagg gatcagcccc gctgtcaca 1752
 ataaagttta ttatgaaaaa aaaaaaaaaa aaaaaaa 1789

310 2
 311 452
 312 FRT
 313 Homo sapiens

400 2
 Met Asp Asp Ile Phe Thr Gln Cys Arg Glu Gly Asn Ala Val Ala Val
 1 5 10 15
 Arg Leu Trp Leu Asp Asn Thr Glu Asn Asp Leu Asn Gln Gly Asp Asp
 20 25 30
 His Gly Phe Ser Pro Leu His Trp Ala Cys Arg Glu Gly Arg Ser Ala
 35 40 45
 Val Val Glu Met Leu Ile Met Arg Gly Ala Arg Ile Asn Val Met Asn
 50 55 60
 Arg Gly Asp Asp Thr Pro Leu His Leu Ala Ala Ser His Gly His Arg
 65 70 75 80
 Asp Ile Val Gln Lys Leu Leu Gln Tyr Lys Ala Asp Ile Asn Ala Val
 85 90 95
 Asn Glu His Gly Asn Val Pro Leu His Tyr Ala Cys Phe Trp Gly Gln
 100 105 110
 Asp Gln Val Ala Glu Asp Leu Val Ala Asn Gly Ala Leu Val Ser Ile
 115 120 125
 Cys Asn Lys Tyr Gly Glu Met Pro Val Asp Lys Ala Lys Ala Pro Leu
 130 135 140
 Arg Glu Leu Leu Arg Glu Arg Ala Glu Lys Met Gly Gln Asn Leu Asn
 145 150 155 160
 Arg Ile Pro Tyr Lys Asp Thr Phe Trp Lys Gly Thr Thr Arg Thr Arg
 165 170 175
 Pro Arg Asn Gly Thr Leu Asn Lys His Ser Gly Ile Asp Phe Lys Gln
 180 185 190
 Leu Asn Phe Leu Thr Lys Leu Asn Glu Asn His Ser Gly Glu Leu Trp
 195 200 205
 Lys Gly Arg Trp Gln Gly Asn Asp Ile Val Val Lys Val Leu Lys Val
 210 215 220
 Arg Asp Trp Ser Thr Arg Lys Ser Arg Asp Phe Asn Glu Glu Cys Pro
 225 230 235 240
 Arg Leu Arg Ile Phe Ser His Pro Asn Val Leu Pro Val Leu Gly Ala
 245 250 255
 Cys Gln Ser Pro Pro Ala Pro His Pro Thr Leu Ile Thr His Trp Met
 260 265 270

Pro Tyr Gly Ser Leu Tyr Asn Val Leu His Glu Gly Thr Asn Phe Val
 275 280 285
 Val Asp Gln Ser Gln Ala Val Lys Phe Ala Leu Asp Met Ala Arg Gly
 290 295 300
 Met Ala Phe Leu His Thr Leu Glu Pro Leu Ile Pro Arg His Ala Leu
 305 310 315 320
 Asn Ser Arg Ser Val Met Ile Asp Glu Asp Met Thr Ala Arg Ile Ser
 325 330 335
 Met Ala Asp Val Lys Phe Ser Phe Gln Cys Pro Gly Arg Met Tyr Ala
 340 345 350
 Pro Ala Trp Val Ala Pro Glu Ala Leu Gln Lys Lys Pro Glu Asp Thr
 355 360 365
 Asn Arg Arg Ser Ala Asp Met Trp Ser Phe Ala Val Leu Leu Trp Glu
 370 375 380
 Leu Val Thr Arg Glu Val Pro Phe Ala Asp Leu Ser Asn Met Glu Ile
 385 390 395 400
 Gly Met Lys Val Ala Leu Glu Gly Leu Arg Pro Thr Ile Pro Pro Gly
 405 410 415
 Ile Ser Pro His Val Cys Lys Leu Met Lys Ile Cys Met Asn Glu Asp
 420 425 430
 Pro Ala Lys Arg Pro Lys Phe Asp Met Ile Val Pro Ile Leu Glu Lys
 435 440 445
 Met Gln Asp Lys
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 #212 - DNA
 #213 - Homo sapiens

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 #211 - 18
 #212 - DNA
 #213 - Homo sapiens

#400 - 4
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 #211 - 19
 #212 - DNA
 #213 - Homo sapiens

#400 - 5
 ccttctccgg ggaactcc 19

#210 - 6
 #211 - 19
 #212 - DNA
 #213 - Homo sapiens

#400 - 6
 cggactccg gctgcagg 19

#210 - 7

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02112- DNA
02113- Homo sapiens

04000-7
gatttatcct cgggactcg 19

02110-8
02111-19
02112- DNA
02113- Homo sapiens

04000-8
gggaaggagg atgaacccc 19

02110-9
02111-19
02112- DNA
02113- Homo sapiens

04000-9
gactgaggac tgtggagtg 19

02110-10
02111-19
02112- DNA
02113- Homo sapiens

04000-10
tggaaagcct gaggactg 19

02110-11
02111-19
02112- DNA
02113- Homo sapiens

04000-11
gagtgaatat gtctccat 19

02110-12
02111-19
02112- DNA
02113- Homo sapiens

04000-12
aacggcgact gcgttgccc 19

02110-13
02111-19
02112- DNA
02113- Homo sapiens

04000-13
tgtttgtccag ccacaggcg 19

02110-14
02111-19
02112- DNA
02113- Homo sapiens

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0210-15	
0211-14	
0212- DNA	
0213- Homo sapiens	
0400-15	
gaagccatga tggcccccc	19
0210-16	
0211-15	
0212- DNA	
0213- Homo sapiens	
0400-16	
caaggccagt gcaaggggg	19
0210-17	
0211-14	
0212- DNA	
0213- Homo sapiens	
0400-17	
cggtccctctt ggcaggccc	19
0210-18	
0211-15	
0212- DNA	
0213- Homo sapiens	
0400-18	
ccacagtaga ggggcccctc	19
0210-19	
0211-16	
0212- DNA	
0213- Homo sapiens	
0400-19	
ccgcatgata aacatctc	18
0210-20	
0211-17	
0212- DNA	
0213- Homo sapiens	
0400-20	
ttgatccgtg cccccccg	18
0210-21	
0211-18	
0212- DNA	
0213- Homo sapiens	
0400-21	
catccccacg gttcattac	19

+210+ 2.
 +211+ 1+
 +212+ LNA
 +213+ Homo sapiens

+400+ 21
 ggaggatca tccccacgg 19

+210+ 1+
 +211+ 1+
 +212+ LNA
 +213+ Homo sapiens

+400+ 23
 ggtgacaga tgcaggggg 19

+210+ 14
 +211+ 1+
 +212+ LNA
 +213+ Homo sapiens

+400+ 24
 atacaaggt gtccatgac 19

+210+ 25
 +211+ 1+
 +212+ LNA
 +213+ Homo sapiens

+400+ 26
 actgcaatag attctgtac 19

+210+ 26
 +211+ 1+
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+400+ 26
 caatgcatg atgtctgac 19

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+210> 31
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+400> 32
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+400> 33
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+211> 19
+212> DNA
+213> Homo sapiens

+400> 34
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+210> 35
+211> 19
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+213> Homo sapiens

+400> 35
tctctgctcc ctctggag 19

+210> 36
+211> 19
+212> DNA
+213> Homo sapiens

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0210 37
0211 19
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0213 Homo sapiens

04008 37
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0210 38
0211 19
0212 DNA
0213 Homo sapiens

04008 38
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0210 39
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04008 39
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0210 40
0211 19
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0213 Homo sapiens

04008 40
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0210 41
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0213 Homo sapiens

04008 41
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19

0210 42
0211 19
0212 DNA
0213 Homo sapiens

04008 42
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0210 43
0211 19
0212 DNA
0213 Homo sapiens

04008 43
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0213> Homo sapiens

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0212> DNA
0213> Homo sapiens

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0213> Homo sapiens

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0210> 47
0211> 19
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0210> 49
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0400> 49
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0211> 19
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19

0210> 51
0211> 19
0212> DNA

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atgagggagca ggtggagac	19
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400 54	
agggtgtgat gagatagg	19
210 55	
211 19	
212 DNA	
213 Homo sapiens	
400 55	
aggatcctat acgcatcc	19
210 56	
211 19	
212 DNA	
213 Homo sapiens	
400 56	
ttcatgtagt acattgtag	19
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211 19	
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400 57	
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211 19	
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+210 - 60
+211 - 19
+212 - RNA
+213 - Homo sapiens

+400 - 19
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+210 - 60
+211 - 19
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+400 - 60
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+210 - 66

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 +400 + 66
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 +400 + 67
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+400+ 81
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0211 10
0212 DNA
0213 Homo sapiens

04009 96
acacacatgt ggcaagtgac

20

0210 97
0211 10
0212 DNA
0213 Homo sapiens

04009 97
ggtgatacct cccatgttgg

20

0210 98
0211 13
0212 DNA
0213 Homo sapiens

04009 98
ggcggggtg atccctccc

19

0210 99
0211 19
0212 DNA
0213 Homo sapiens

04009 99
taataaaatt tattgtgac

19